



SEQUENCE LISTING

5 <110> Tillett, D
Thomas, T

<120> A method of sequestering and/or purifying a polypeptide

10 <130> nuc2004

<140> 10/785,452

<141> 2004-02-25

15 <150> PCT/AU02/01159

<151> 2002-08-27

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<211> 714

25 <212> DNA

<213> Aequorea victoria

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Met Ser Lys Gly Glu Glu Leu Phe Thr Gly Val Val Pro Ile Leu
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35 gtt gaa tta gat ggc gat gtt aat ggg caa aaa ttc tct gtc agt
90
Val Glu Leu Asp Gly Asp Val Asn Gly Gln Lys Phe Ser Val Ser
20 25 30

40 gga gag ggt gaa ggt gat gca aca tac gga aaa ctt acc ctt aaa
135
Gly Glu Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys
35 40 45

45 ttt att tgc act act ggg aag cta cct gtt cca tgg cca aca ctt
180
Phe Ile Cys Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu
50 55 60

50

50

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ccc aac gaa aag aga gat cac atg atc ctt ctt gag ttt gta aca
675
Pro Asn Glu Lys Arg Asp His Met Ile Leu Leu Glu Phe Val Thr
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gct gct ggg att aca cat ggc atg gat gaa cta tac aaa 714
Ala Ala Gly Ile Thr His Gly Met Asp Glu Leu Tyr Lys
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<213> Escherichia coli

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25
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90
Ile Gly Val Gly Gly Gly Gly Gly Asn Ala Val Glu His Met Val
20 25 30

30
cgc gag cgc att gaa ggt gtt gaa ttc ttc gcg gta aat acc gat
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Arg Glu Arg Ile Glu Gly Val Glu Phe Phe Ala Val Asn Thr Asp
35 40 45

35
gca caa gcg ctg cgt aaa aca gcg gtt gga cag acg att caa atc
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Ala Gln Ala Leu Arg Lys Thr Ala Val Gly Gln Thr Ile Gln Ile
50 55 60

40
ggt agc ggt atc acc aaa gga ctg ggc gct ggc gct aat cca gaa
225
Gly Ser Gly Ile Thr Lys Gly Leu Gly Ala Gly Ala Asn Pro Glu
65 70 75

45
gtt ggc cgc aat gcg gct gat gag gat cgc gat gca ttg cgt gcg
270
Val Gly Arg Asn Ala Ala Asp Glu Asp Arg Asp Ala Leu Arg Ala
80 85 90

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	gaa atg gct atc tct tct ccg ctg ctg gaa gat atc gac ctg tct	
	765	
	Glu Met Ala Ile Ser Ser Pro Leu Leu Glu Asp Ile Asp Leu Ser	
	245	250 255
5	ggc gcg cgc ggc gtg ctg gtt aac atc acg gcg ggc ttc gac ctg	
	810	
	Gly Ala Arg Gly Val Leu Val Asn Ile Thr Ala Gly Phe Asp Leu	
	260	265 270
10	cgt ctg gat gag ttc gaa acg gta ggt aac acc atc cgt gca ttt	
	855	
	Arg Leu Asp Glu Phe Glu Thr Val Gly Asn Thr Ile Arg Ala Phe	
	275	280 285
15	gct tcc gac aac gcg act gtg gtt atc ggt act tct ctt gac ccg	
	900	
	Ala Ser Asp Asn Ala Thr Val Val Ile Gly Thr Ser Leu Asp Pro	
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20	gat atg aat gac gag ctg cgc gta acc gtt gtt gcg aca ggt atc	
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	Asp Met Asn Asp Glu Leu Arg Val Thr Val Val Ala Thr Gly Ile	
	305	310 315
25	ggc atg gac aaa cgt cct gaa atc act ctg gtg acc aat aag cag	
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<213> Human rhinovirus

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15 atg act ata aca acc tca aag gga gag ttc aca ggg tta ggc ata
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     20      25      30

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    His Asp Arg Val Cys Val Ile Pro Thr His Ala Gln Pro Gly Asp
     35      40      45

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    Asp Val Leu Val Asn Gly Gln Lys Ile Arg Val Lys Asp Lys Tyr
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    Lys Leu Val Asp Pro Glu Asn Ile Asn Leu Glu Leu Thr Val Leu
     65      70      75

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   270
    Thr Leu Asp Arg Asn Glu Lys Phe Arg Asp Ile Arg Gly Phe Ile
     80      85      90

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    Ser Glu Asp Leu Glu Gly Val Asp Ala Thr Leu Val Val His Ser
     95     100     105

45 aat aac ttt acc aac act atc tta gaa gtt ggc cct gta aca atg
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    Asn Asn Phe Thr Asn Thr Ile Leu Glu Val Gly Pro Val Thr Met
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50 gca gga ctt att aat ttg agt agc acc ccc act aac aga atg att
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cgt	tat	gat	tat	gca	aca	aaa	act	ggg	cag	tgt	gga	ggt	gtg	ctg
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Arg	Tyr	Asp	Tyr	Ala	Thr	Lys	Thr	Gly	Gln	Cys	Gly	Gly	Val	Leu
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tgt	gct	act	ggt	aag	atc	ttt	ggt	att	cat	gtt	ggc	ggt	aat	gga
495														
Cys	Ala	Thr	Gly	Lys	Ile	Phe	Gly	Ile	His	Val	Gly	Gly	Asn	Gly
				155					160					165

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aga	caa	gga	ttt	tca	gct	caa	ctt	aaa	aaa	caa	tat	ttt	gta	gag
540														
Arg	Gln	Gly	Phe	Ser	Ala	Gln	Leu	Lys	Lys	Gln	Tyr	Phe	Val	Glu
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aaa	caa	546												
Lys	Gln													
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